

AMENDMENTS TO THE CLAIMS

Claim 1 (currently amended) A method comprising the step of:

inhibiting attachment of *C. albicans* to human tissue by interfering with the transcription of hyphal specific genes in *C. albicans* mediated by *eis* acting sequences, wherein cell growth of said *C. albicans* is inhibited DNA binding proteins specific to UAS regions of the promoter of *HWP1* present during germ tube formation in said *C. albicans*.

Claims 2-20 (canceled)

Claim 21 (currently amended) The method of claim 20 ~~1~~, wherein said ~~*HWP1*~~ promoter of *HWP1* region comprises the following isolated DNA sequence [SEQ. ID. NO:1]:

GGATCTTTCTTTTTCATTTCCCTTAAAACCGATCAAGAAAGAAAGTGGAAATAAA
GCTATGATAAATGTTGATTTTGTGTAATTCAATCAACTAAGCACGTTTGACAGTT
AAAAAGTACGTTGTTGTTGTCCTCGTCTCGTCTAATTTCTGTTGACGAGGATTAAT
AACAAGAAATACAGGAAACCTCCAAAAAATAATTTTGGACCTTACACGCACA
TAAATTGCGGATAAACTTGCCATAATAAAACTCTTTGAAACATACGATATGTTA
TTCTTTTTCATAACTGGAATATTTTGTCTTTTTTTTAACATTATGAACAATTGAAAA
AAAAAGGAAATGAAAAGGTAAGAGTTGCCTAACCATTGAAAATAATAGGCTAAG
GTTTTTCCTGATGCGTTTAACTAAAAAGGAAATAACAAAAGTTATTAGCGATAAC
CTGCGTAAGGTGTCAACAAAATATATTTTGCACGTTAGCTCTATAGAAAATATAC
AACTAAATCCTTAAGGAATTTCTCTATATATAATAGGAAATCCCTCTCACAGT
GAACTGAATTATCCATCTGAATTATCAGTCCACTAATTCCATCAATAAAATAGAT
TAGTGTATTGTTCTCTTCAGTACAATTACTACCATTATGCAATGCTAGCTTATTGT
TCATAATTAGCCATGTTGCACACCCTAATTGCAACATTAAGTATCCATATTTTT
CTTGTCCTTCTTTGTTTTTTTCTAACAAAAATGTTCCAGAATTTTTTAAAAAATATT
TGAAAAAACACATAACACTTTGAGTATGATAATCAACTATTGACTTGTTTTGA
AAGTAAAGAATCAAATTTTTTTCTAACTCGACTAATGCACTTTACATCAACTGGA
TGTTATTTGCATCTACTACTATAAGCTCAAACAAATTATCTTTCAAAAATGTTATA
ATTAACAAGTCATCTATAATTCTTTGGATCCAAAAACAAGGAATTTCGGAAATTCT
GACGATAAATGTCGACTCACAATTCATTGTAAAAAGGGAGAGTTTTGGTAGGCTC
ATAATCGCTTATAATGTACCTCTAAAGTAATCTAAAACAAACACAACCTTTCTAA
AACCTATAATAATAACCCTAATGGCTCACAACCGGGATAATGTTAGTTAGCCCAG
CTGTTTTTTTTTTGCCTTATTTTTATGACTACATTTTGTTCACTTTTTGTGCGACT
TTAATACCGTTTTTGAACCTTCTCTTTGTATCACCTGTATCCGCCTTTTTTAACATA

GCAACTCTTGTAAGTCCCTTTCTTTTCCCACTATTTTATCATTCTTGAAATATGT
AATCAGAATAGTTTTTCAAAAACATAAAATAACGGTCAAAATAACCGGCTATTTT
CAATTTCCATTCAACTTGTTTTCTCAACAATATCAAACACAACAGGAATCTCCTAT
AGTCACTCGCTTTTAGTTTCGTCAATATG;

~~including any insertions, deletions, mutations, or modifications.~~

Claims 22-27 (canceled)

Claim 28 (currently amended) The method of claim 4~~1~~, wherein said ~~cis-regulatory~~
~~elements~~ UAS regions comprise a NIT2 binding site.

Claim 29 (currently amended) The method of claim 7~~28~~, wherein said DNA BP is encoded
by a nucleotide sequence for the DNA binding domain that is homologous to a nucleotide
sequence encoding the DNA binding domain of NIT2 binding proteins.

Claim 30 (original) The method of claim 29, wherein the DNA BP is selected from the group
consisting of GAT99, GAT-1, or GATA-like binding proteins.

Claim 31 (original) The method of claim 29, wherein said NIT2 DNA binding domain
comprises the protein sequence [SEQ. ID. NO: 2]:

CTNCFTQTTPLWRRNPDGQPLCNACGLFLKLHGVRPLSLKTDVIKKRNR.

Claim 32 (original) The method of claim 29, wherein the DNA binding domain of said DNA
BP comprises the protein sequence [SEQ. ID. NO: 3]:

CTNCGTKTTPLWRRNPQGQPLCNACGLFLKLHGVRPLSLKTDVIKKRQR.

Claims 33-43 (canceled)